Appendix I: Alignment of SEQ ID NO: 17 (CRF, 10/20/2009) and I48378

```
Appendix B
<!--StartFragment-->RESULT 1
148378
hairless protein - mouse
C; Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #secuence_revision 02-Jul-1996 #text_change 09-Jul-2004
C;Accession: I48378
R; Cachon-Gonzalez, M.B.; Fenner, S.; Coffin, J.M.; Moran, C.; Best, S.; Stoye, J.P.
Proc. Natl. Acad. Sci. U.S.A. 91, 7717-7721, 1994
A; Title: Structure and expression of the hairless gene of mice.
A:Reference number: I48378; MUID:94329587; PMID:8052649
A:Accession: I48378
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: mRNA
A; Residues: 1-1182 < RES>
A; Cross-references: UNIPROT: Q61645; UNIPARC: UPI0000028C24; EMBL: Z32675; NID: q531706; P
 Query Match
                     99.4%; Score 3682; DB 2; Length 1182;
 Best Local Similarity
                    99.4%;
 Matches 689; Conservative
                          1; Mismatches
                                        3; Indels
                                                  0; Gaps
                                                              0:
Qу
          1 VTOCOSCVOAAGEVGVLTGHSOKSRRSPLEEKOLEEEDSSATSEEGGGGPGPEASLNKGL 60
           Db
        490 VTOCOSCVCAAGEVGVLTGHSOKSRRSPLEEKOLEEEDSSATSEEGGGGPGPEASLNKGL 549
        61 AKHLLSGLGDRLCRLLRKEREALAWAQREGQGPAMTEDSPGIPHCCSRCHHGLFNTHWRC 120
Qу
           Db
        550 AKHLLSGLGDRLCRLLRKEREALAWAQREGQGPAMTEDSPGIPHCCSRCHHGLFNTHWRC 609
        121 SHCSHRLCVACGRIAGAGKNREKTGSQEQHTDDCAQEAGHAACSLILTQFVSSQALAELS 180
Qy
           Dh
        610 SHCSHRLCVACGRIAGAGKNREKTGSQEQHTDDCAQEAGHAACSLILTQFVSSQALAELS 669
        181 TVMHCAWARFDIRGHCFCOVDARVWAPGDGGOOKEPTEKTPPTPOPSCNGDSNRTKDIKE 240
Qv
           670 TVMHOWWAKFDIRGHCFCQVDARVWAPGDGGQQKEPTEKTPPTPQPSCNGDSNRTKDIKE 729
Db
        241 ETPDSTESPAEDGAGRSPLPCPSLCELLASTAVKLCLGHDRIHMAFAPVTPALPSDDRIT 300
Qу
           730 ETPDSTESPAEDGAGRSPLPCPSLCELLASTAVKLCLGHDRIHMAFAPVTPALPSDDRIT 789
Dh
       301 NILDSIIAQVVERKIQEKALGPGLRAGSGLRKGLSLPLSPVRTRLSPPGALLWLQEPRPK 360
Qу
           nh
        790 NILDSIIAQVVERKIQEKALGPGLRAGSGLRKGLSLPLSPVRTRLSPPGALLWLQEPRPK 849
       361 HGFHLFQEHWRQGQPVLVSGIQKTLRLSLWGMEALGTLGGQVQSLTALGPPQPTNLDSTA 420
Qy
           850 HGFHLFQEHWRQGQPVLVSGIQKTLRLSLWGMEALGTLGGQVQTLTALGPPOPTNLDSTA 909
Dh
        421 FWEGFSHPETRPKLDEGSVLLLHRTLGDKDASRVQMLVSLPLPEYCAHQGKLNLASYLP 480
Qν
           Dh
       910 FWEGFSHPETRPKLDEGSVLLLHRTLGDKDASRVONLASSLPLPEYCAHOGKLNLASYLP 969
Qv
       481 LGLTLHPLEPQLWAAYGVNSHRGHLGTKNLCVEVSDLISILVHAEAQLPPWYRAQKDFLS 540
           Db
       970 LGLTLEPLEPQLWAAYGVNSHRGHLGTKNLCVEVSDLISILVHAEAQLPPWYRAQKDFLS 1029
       541 GLDGEGLWSPGSQTSTVWHVFRAQDAQRIRRFLQMVCPAGAGTLEPGAPGSCYLDAGLRR 600
Qу
           Db
       1030 GLDGEGLWSPGSQTSTVWHVFRAQDAQRIRRFLQMVCPAGAGTLEPGAPGSCYLDAGLRR 1089
       601 RLREEWGVSCWTLLQAPGEAVLVPAGAPHQVQGLVSTISVTQHFLSPETSALSAQLCHQG 660
Qν
```

Qy 661 ASLPPDHRMLYAQMDRAVFQAVKAAVGALQEAK 693

Db 1150 ASLPPDHRMLYAQMDRAVFQAVKAAVGALQEAK 1182

<!--EndFragment-->

Db